

PCT

RAW SEQUENCE LISTING

DATE: 11/04/2003 TIME: 13:48:52

PATENT APPLICATION: US/09/601,644A

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Input Set : A:\seqlist.txt

Output Set: N:\CRF4\11042003\I601644A.raw

- 3 <110> APPLICANT: Gareipy, Jean
- 4 Bray, Mark Robert
- 6 <120> TITLE OF INVENTION: Cytotoxic Heteromeric Protein Combinatorial Libraries
- 8 <130> FILE REFERENCE: MMC.P-001
- 10 <140> CURRENT APPLICATION NUMBER: 09/601,644A
- C--> 11 <141> CURRENT FILING DATE: 2003-12-27
  - 13 <150> PRIOR APPLICATION NUMBER: PCT/CA98/01137
  - 14 <151> PRIOR FILING DATE: 1998-12-08
  - 16 <160> NUMBER OF SEQ ID NOS: 4
  - 18 <170> SOFTWARE: PatentIn version 3.2

Does Not Comply
Corrected Disketto Needed

## ERRORED SEQUENCES

- E--> 111 <210> SEQ ID NO: 65 3
  - 113 <212> TYPE: DNA
  - 114 <213> ORGANISM: artificial
  - 116 <220> FEATURE:
  - 117 <223> OTHER INFORMATION: Primer
  - 119 <220> FEATURE:
  - 120 <221> NAME/KEY: misc\_feature
  - 121 <222> LOCATION: (25)..(26) ...
  - 122 <223> OTHER INFORMATION: n 's a, c, g, or t
  - 124 <220> FEATURE:
  - 125 <221> NAME/KEY: misc\_feature /
  - 126 <222> LOCATION: (28)..(29)
  - 127 <223> OTHER INFORMATION: n is a, c, g, or t
  - 129 <220> FEATURE:
  - 130 <221> NAME/KEY: misc\_feature/
  - 131 <222> LOCATION: (31)..(32)
  - 132 <223> OTHER INFORMATION: n is a, c, g, or t
  - 134 <220> FEATURE:
  - 135 <221> NAME/KEY: misc feature
  - 136 <222> LOCATION: (34)..(35)
  - 137 <223> OTHER INFORMATION: n is a, c, g, or t
  - 139 <220> FEATURE:
  - 140 <221> NAME/KEY: misc feature
  - 141 <222> LOCATION: (37)..(38)
  - 142 <223> OTHER INFORMATION: n is a, c, q, or t
  - 144 <220> FEATURE:
  - 145 <221> NAME/KEY: misc feature
  - 146 <222> LOCATION: (67)..(68)

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147 <223> OTHER INFORMATION: n is a, c, g, or t 149 <220> FEATURE: 150 <221> NAME/KEY: misc\_feature 151 <222> LOCATION: (70)..(71) 152 <223> OTHER INFORMATION: n is a, c, g, or t 154 <220> FEATURE: 155 <221> NAME/KEY: misc feature 156 <222> LOCATION: (73)..(74) 157 <223> OTHER INFORMATION: n is a, c, g, or t 159 <220> FEATURE: 160 <221> NAME/KEY: misc\_feature 161 <222> LOCATION: (76)..(77) 162 <223> OTHER INFORMATION: n is a, c, g, or t E--> 164 <400> SEQUENCE: 3 W--> 165 aaggtggagt atacaaaata taatnnsnns nnsnnsnnsa cagttaaagt gggtgataaa 60 167 gaattannsn nsnnsnnstg gaatetteag tete 94

E--> 170 <210> SEQ ID NO: 4

## VERIFICATION SUMMARY

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Input Set : A:\seqlist.txt

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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:111 M:214 E: (33) Seq.# missing, SEQ ID NO:3

L:164 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:4 differs:3

L:165 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0

M:341 Repeated in SeqNo=4

L:170 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO:4